



SEQUENCE LISTING

<110> Goedegebuur, Frits
Gualfetti, Peter
Mitchinson, Colin
Neefe, Paulien

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<150> US 60/456,368

<151> 2003-03-21

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<213> Hyprocrea jecorina

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<212> PRT

<213> Hyprocrea jecorina

<400> 2

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 <213> *Hyprocrea orientalis*

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<212> DNA

<213> *Hyprocrea schweintzii*

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<213> *Hyprocrea schweintzii*

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<211> 497

<212> PRT

<213> *Hyprocrea schweintzii*

<400> 8

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Val Ile Asp Ala Asn Trp Arg Trp Thr His Ala Thr Asn Ser Ser Thr
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Asn Cys Tyr Asp Gly Asn Thr Trp Ser Ser Thr Leu Cys Pro Asp Asn
      50             55             60
Glu Thr Cys Ala Lys Asn Cys Cys Leu Asp Gly Ala Ala Tyr Ala Ser
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<213> Trichoderma konilangbra

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<210> 11

<211> 498

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<213> Trichoderma konilangbra

<220>

<221> VARIANT

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20

25

30

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35

40

45

Asn Cys Tyr Asp Gly Asn Thr Trp Ser Ser Ser Leu Cys Pro Asp Asn

50

55

60

Glu Ser Cys Ala Lys Asn Cys Cys Leu Asp Gly Ala Ala Tyr Ala Ser

65

70

75

80

Thr Tyr Gly Val Thr Thr Ser Ala Asp Ser Leu Ser Ile Gly Phe Val

85

90

95

Thr Gln Ser Gln Gln Lys Asn Val Gly Ala Arg Leu Tyr Leu Met Ala

100

105

110

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115

120

125

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Pro	Arg	Asp	Leu	Lys	Phe	Ile	Asn	Gly	Glu	Ala	Asn	Val	Glu	Gly	Trp
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Cys Leu

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 <213> *Trichoderma pseudokoningii*

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 <213> Trichoderma pseudokoningii

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 Ala

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 <212> PRT
 <213> Trichoderma pseudokoningii

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 35 40 45
 Asn Cys Tyr Asp Gly Asn Thr Trp Ser Ser Thr Leu Cys Pro Asp Asn
 50 55 60
 Glu Thr Cys Ala Lys Asn Cys Cys Leu Asp Gly Ala Ala Tyr Ala Ser
 65 70 75 80
 Thr Tyr Gly Val Thr Thr Ser Ala Asp Ser Leu Ser Ile Gly Phe Val
 85 90 95
 Thr Gln Ser Ala Gln Lys Asn Val Gly Ala Arg Leu Tyr Leu Met Ala
 100 105 110
 Ser Asp Thr Thr Tyr Gln Glu Phe Thr Leu Leu Gly Asn Glu Phe Ser
 115 120 125
 Phe Asp Val Asp Val Ser Gln Leu Pro Cys Gly Leu Asn Gly Ala Leu
 130 135 140
 Tyr Phe Val Ser Met Asp Ala Asp Gly Gly Val Ser Lys Tyr Pro Thr
 145 150 155 160
 Asn Thr Ala Gly Ala Lys Tyr Gly Thr Gly Tyr Cys Asp Ser Gln Cys
 165 170 175
 Pro Arg Asp Leu Lys Phe Ile Asn Gly Glu Ala Asn Val Glu Gly Trp
 180 185 190

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Cys	Cys	Ser	Glu	Met	Asp	Ile	Trp	Glu	Ala	Asn	Ser	Ile	Ser	Glu	Ala
	210					215					220				
Leu	Thr	Pro	His	Pro	Cys	Thr	Thr	Val	Gly	Gln	Glu	Ile	Cys	Asp	Gly
225					230					235					240
Asp	Ser	Cys	Gly	Gly	Thr	Tyr	Ser	Gly	Asp	Arg	Tyr	Gly	Gly	Thr	Cys
				245					250					255	
Asp	Pro	Asp	Gly	Cys	Asp	Trp	Asn	Pro	Tyr	Arg	Leu	Gly	Asn	Thr	Ser
			260					265					270		
Phe	Tyr	Gly	Pro	Gly	Ser	Ser	Phe	Ala	Leu	Asp	Thr	Thr	Lys	Lys	Leu
		275					280					285			
Thr	Val	Val	Thr	Gln	Phe	Glu	Thr	Ser	Gly	Ala	Ile	Asn	Arg	Tyr	Tyr
	290					295					300				
Val	Gln	Asn	Gly	Val	Thr	Phe	Gln	Gln	Pro	Asn	Ala	Glu	Leu	Gly	Ser
305					310					315					320
Tyr	Ser	Gly	Asn	Ser	Leu	Asp	Asp	Asp	Tyr	Cys	Ala	Ala	Glu	Glu	Ala
				325					330					335	
Glu	Phe	Gly	Gly	Ser	Ser	Phe	Ser	Asp	Lys	Gly	Gly	Leu	Thr	Gln	Phe
			340					345					350		
Lys	Lys	Ala	Thr	Ser	Gly	Gly	Met	Val	Leu	Val	Met	Ser	Leu	Trp	Asp
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Asp	Tyr	Tyr	Ala	Asn	Met	Leu	Trp	Leu	Asp	Ser	Thr	Tyr	Pro	Thr	Asn
	370				375						380				
Glu	Thr	Ser	Ser	Thr	Pro	Gly	Ala	Val	Arg	Gly	Ser	Cys	Ser	Thr	Ser
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Ser	Gly	Val	Pro	Ala	Gln	Leu	Glu	Ser	Gln	Ser	Ser	Asn	Ala	Lys	Val
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Val	Tyr	Ser	Asn	Ile	Lys	Phe	Gly	Pro	Ile	Gly	Ser	Thr	Gly	Asn	Ser
			420					425					430		
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		435				440						445			
Thr	Arg	Arg	Pro	Ala	Thr	Ser	Thr	Gly	Ser	Ser	Pro	Gly	Pro	Thr	Gln
	450				455						460				
Thr	His	Tyr	Gly	Gln	Cys	Gly	Gly	Ile	Gly	Tyr	Ser	Gly	Pro	Thr	Val
465					470					475					480
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		20						25					30		
Val	Ile	Asp	Ala	Asn	Trp	Arg	Trp	Thr	His	Ala	Thr	Asn	Ser	Ser	Thr
		35				40						45			
Asn	Cys	Tyr	Asp	Gly	Asn	Thr	Trp	Ser	Ser	Thr	Leu	Cys	Pro	Asp	Asn
	50				55						60				
Glu	Thr	Cys	Ala	Lys	Asn	Cys	Cys	Leu	Asp	Gly	Ala	Ala	Tyr	Ala	Ser
65				70						75				80	
Thr	Tyr	Gly	Val	Thr	Thr	Ser	Gly	Asn	Ser	Leu	Ser	Ile	Gly	Phe	Val
				85					90					95	
Thr	Gln	Ser	Ala	Gln	Lys	Asn	Val	Gly	Ala	Arg	Leu	Tyr	Leu	Met	Ala
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19